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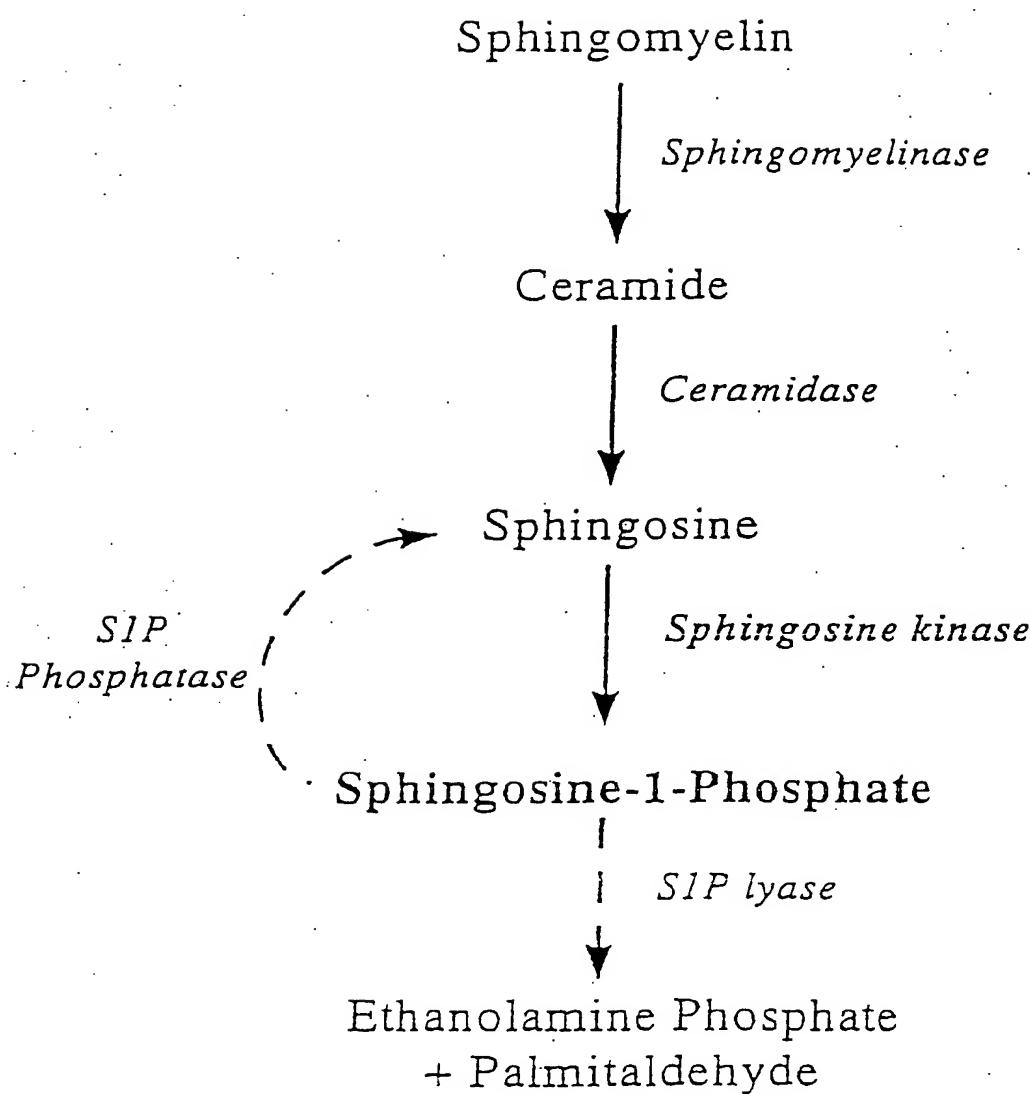


Figure 1

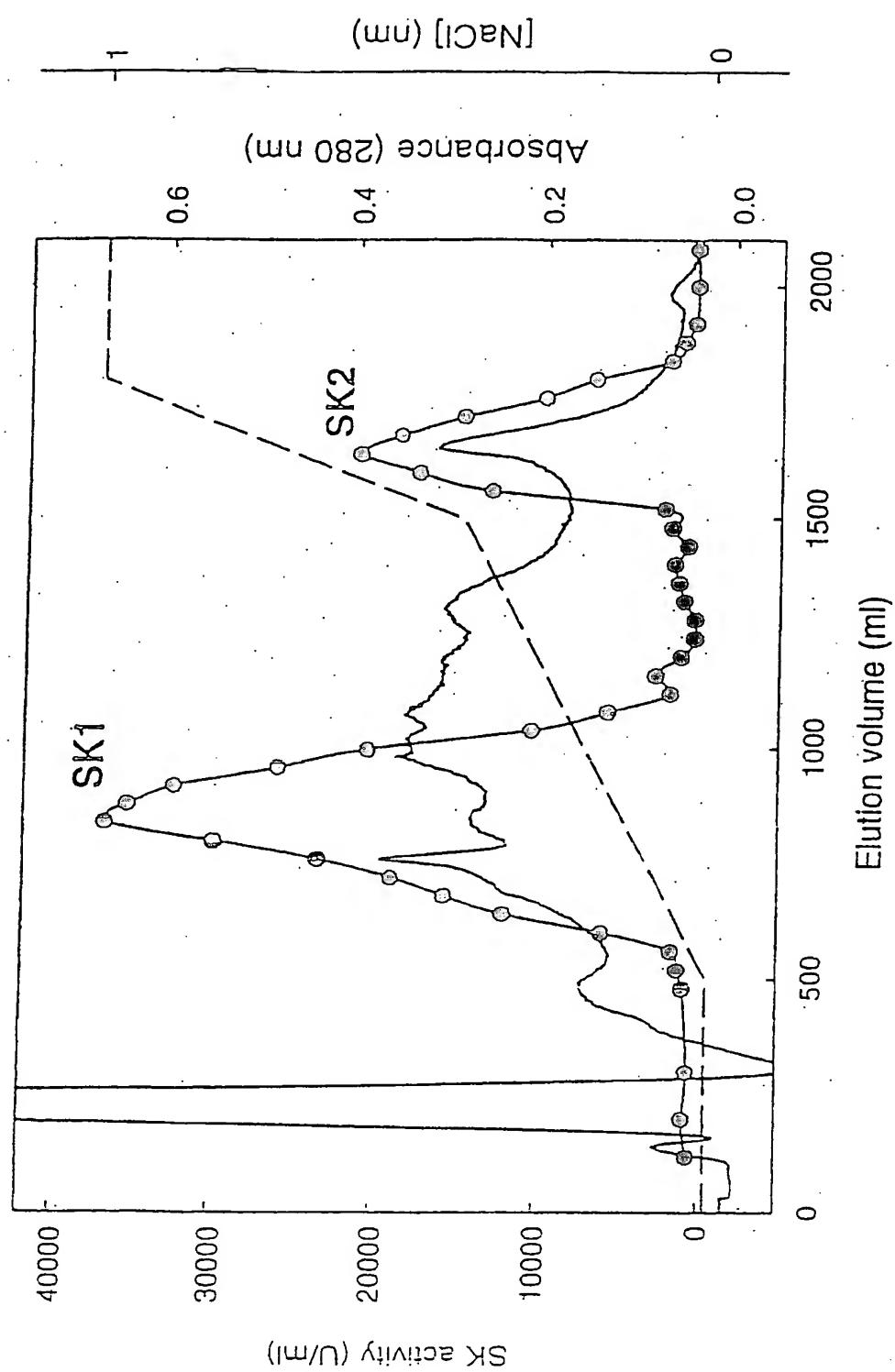


Figure 2

Figure 3A

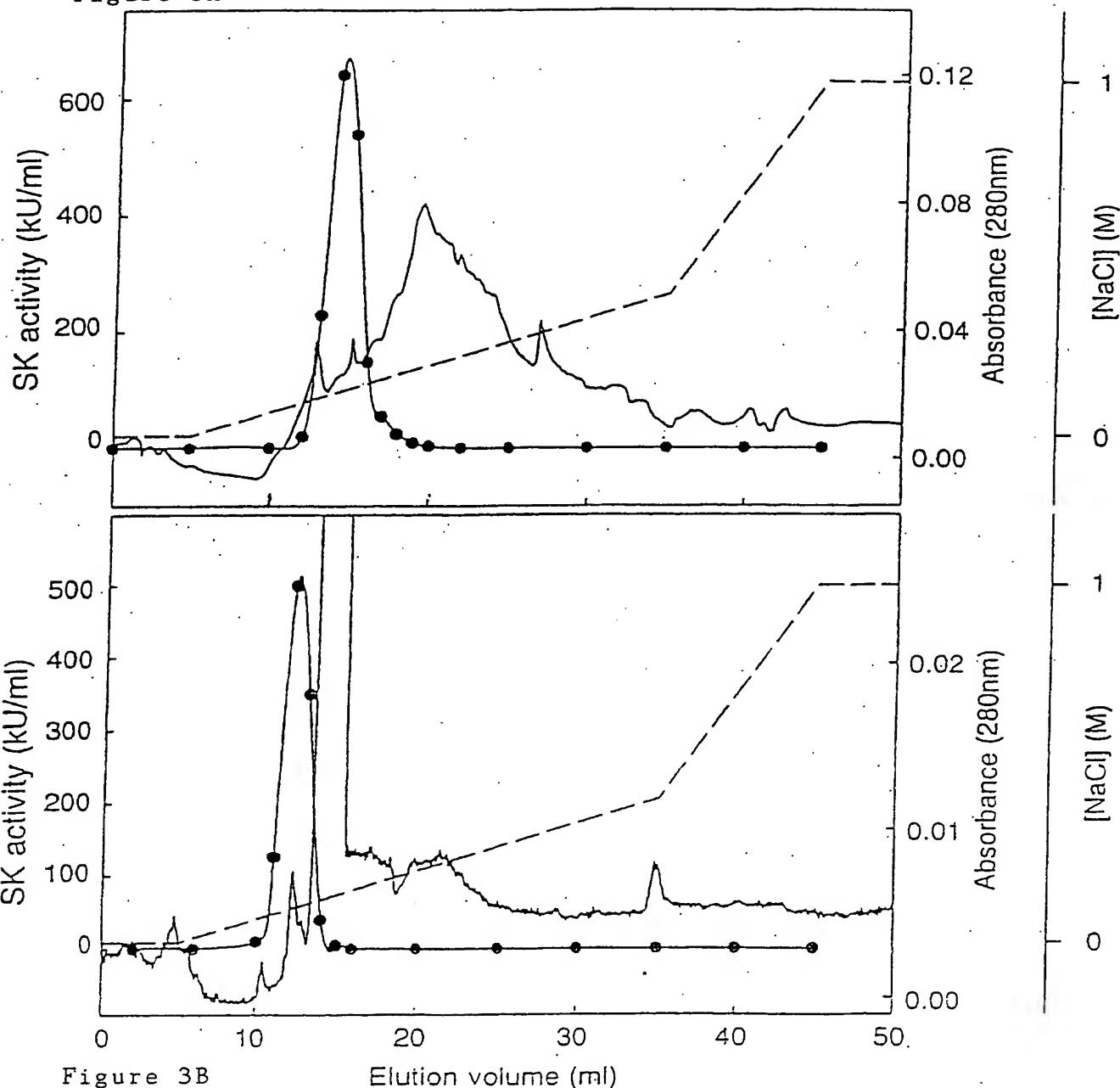


Figure 3B

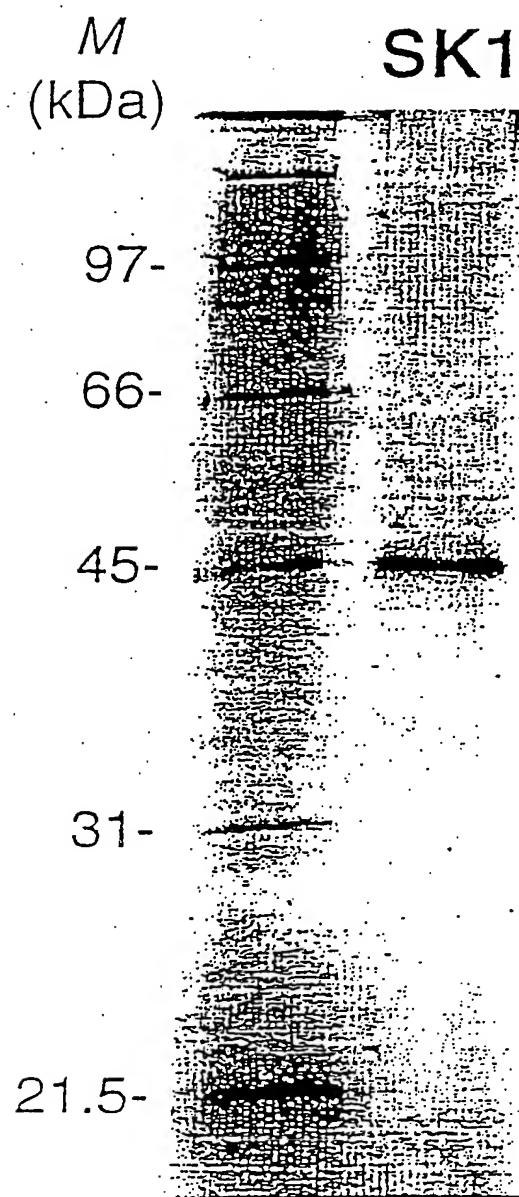


Figure 4

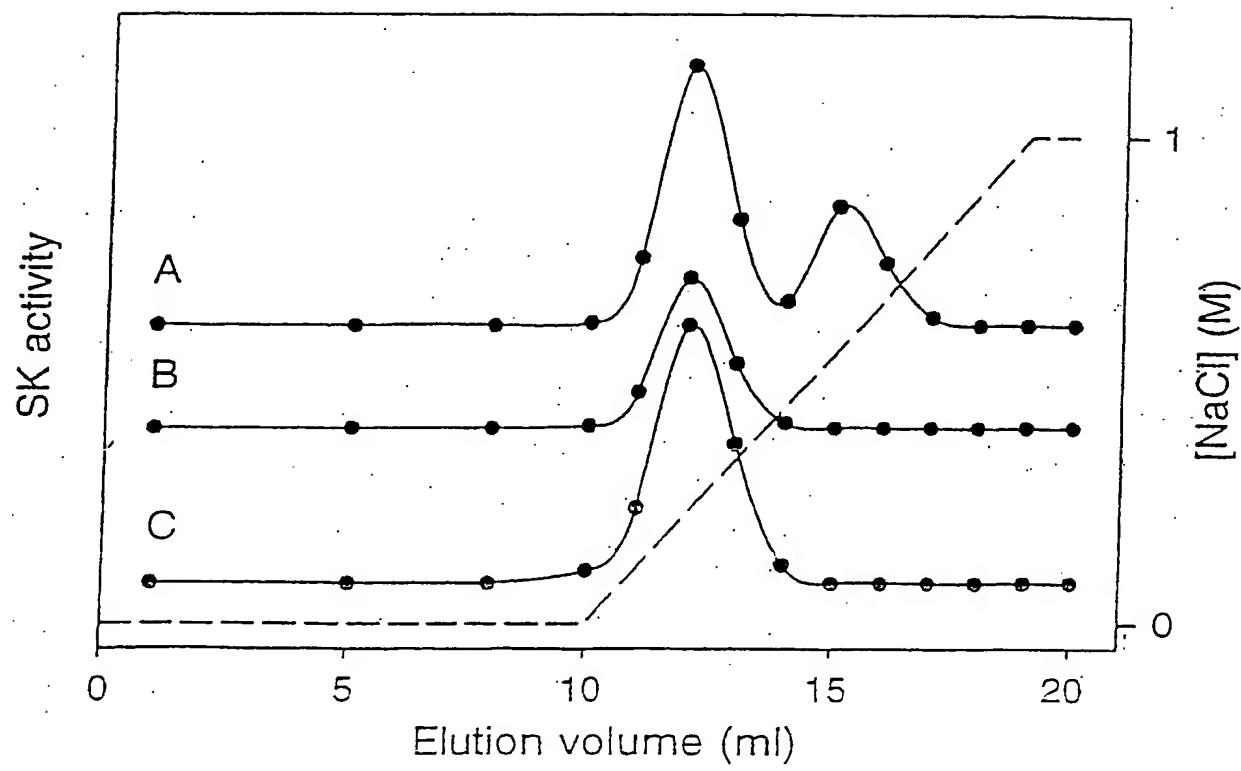


Figure 5

Cloning of HUVEC SK

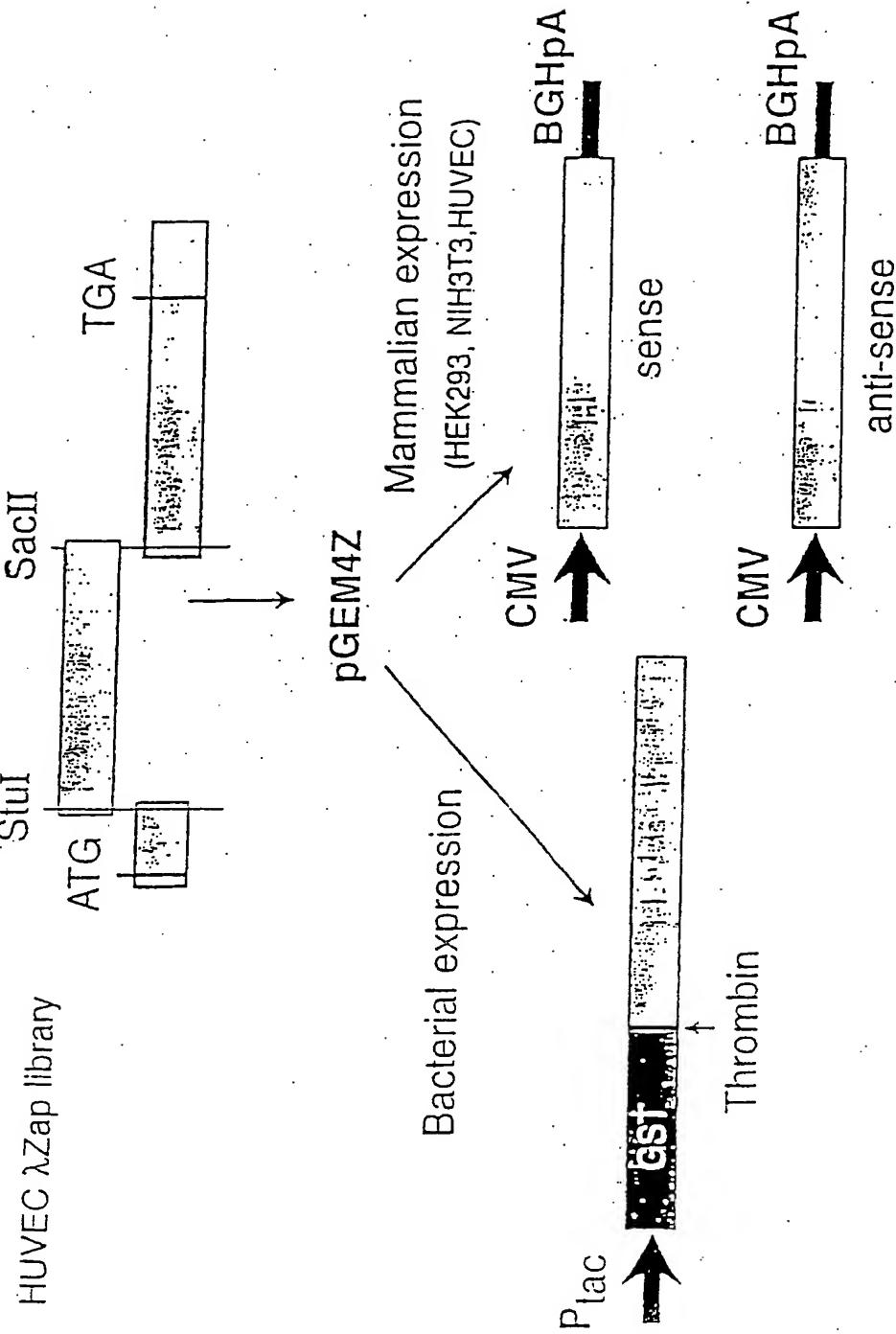


Figure 6

Figure 7A-1

195 G T F L R A A L R T Y R G R L A Y L P V G R V G 219
 601 GCACTTCTGGCTCTGGCAACTTACCGGGCGACTGGCTTACCTCCCTGTAGGAAGACTGGGT 675
 220 S K T P A S P V V Q Q G P V D A H L V P L E E P 244
 676 TCCAGACACCTGGCTCCCCGGTGTGGTCCAGCAGGGCCGGTAGATGCACACCTTGTGCCACTGGAGGCCA 750
 245 V P S H W T V V P D E D F V L V L A L L H S H L G 269
 751 GTGCCCTCACTGGACAGTGGACTGGCCGACGGAGACTTTGTGCTAGGCTGGACTGTGCACTGGCACCTGGCC 825
 270 S E N F A A P M G R C A A G V M H L F Y V R A G V 294
 826 AGTGAGATGTTGCTGCACCCATGGCCGCTGTGCAAGCTGGGTCAATGCATCTGTTCTACGTGGAGTGC 900
 295 S R A M L L R L F L A M E K G R H M E Y E C P Y L 319
 901 TCTCGTCATGCTGCTGGCCTCTGGCCATGGAGAAGGGCAGGCATATGGAGTATGAATGCCCTACTTG 975
 320 V Y V P V A F R L E P K D G K G M F A V D G E L 344
 976 GTATATGCCCCGGTGGTGCCTTCCGCTTGGAGCCCCAAGGATGGAAAGGTATGTTGCAAGTGGATGGGAATTG 1050
 345 M V S E A V Q G Q V H P N Y F W M V S G C V E P P 369
 1051 ATGGTTAGCGAGGCCGTGCAGGGCCAGGTGCACCCAAACTACTTCTGGATGGTCAGGGTTGGCTGGAGCCCCCG 1125
 370 P S W K P Q Q M P P E E P L 384
 1126 CCCAGCTGGAAGCCCCAGCAGATGCCACCGCCAGMAAGGCCCTTATga 1173

Figure 7A-2

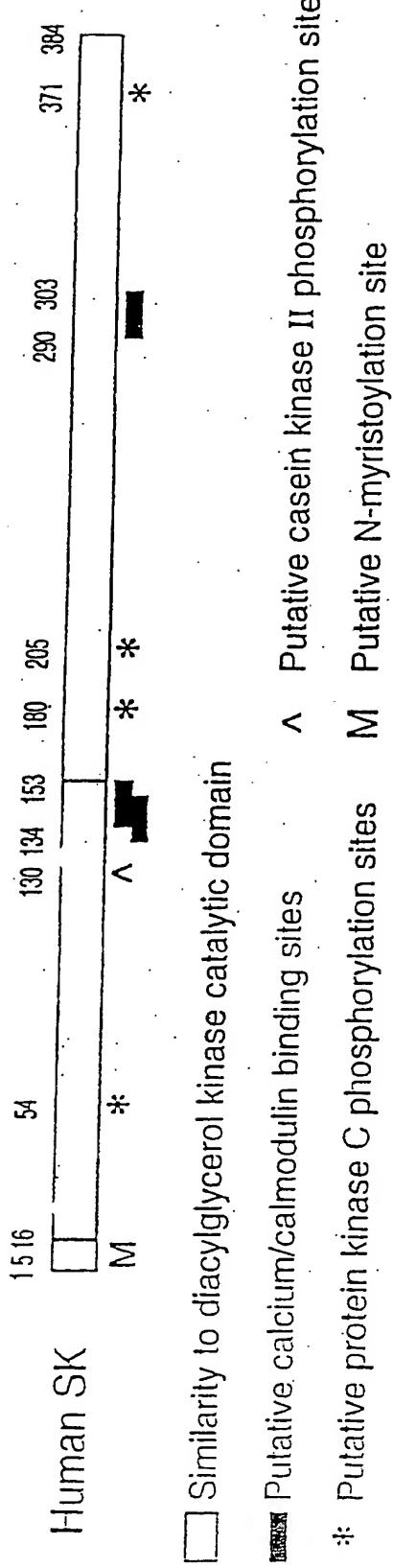


Figure 7B

Figure 8A

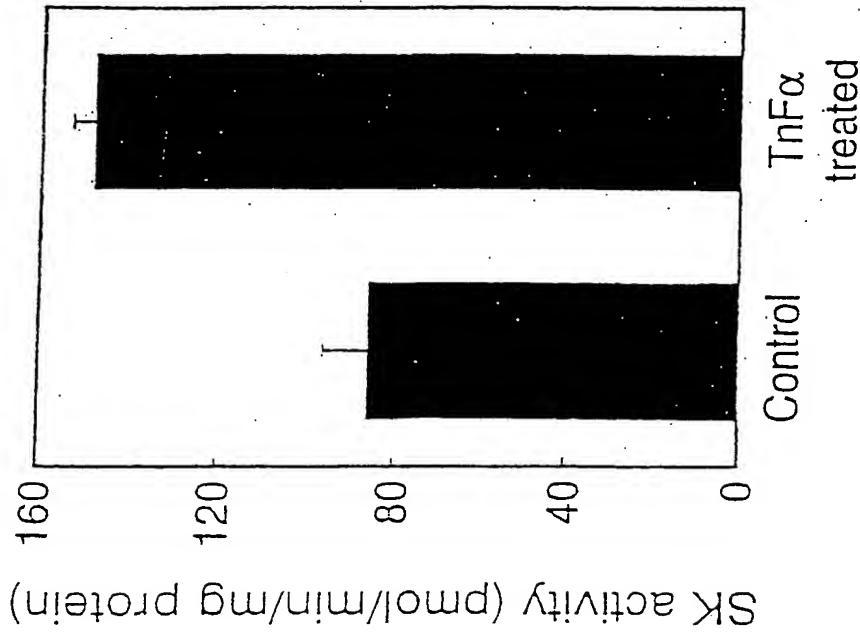
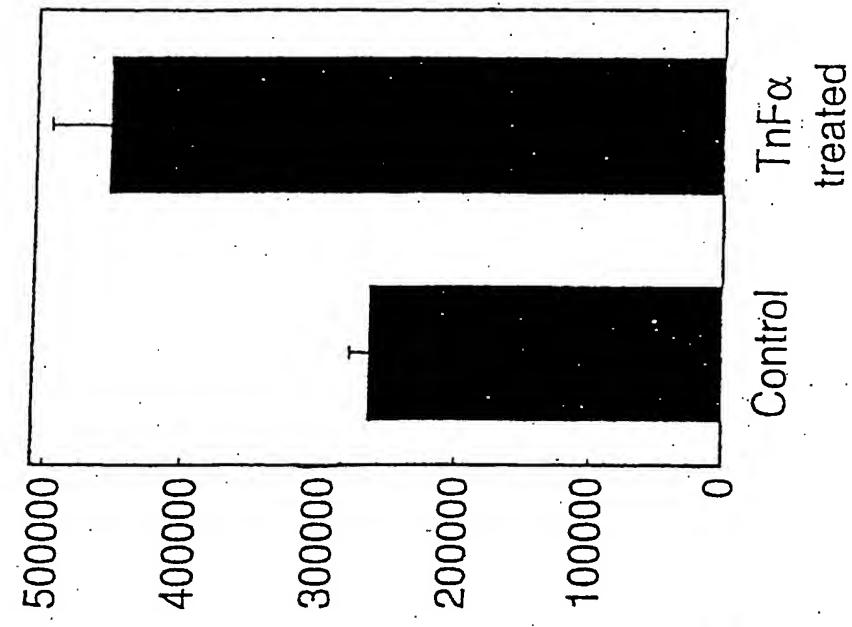


Figure 8B



hSK	1	MDPAGGPRGVLP PR PCRVL W LNPRGGKGKALQLE FRSHVQ Q FL LAEEAIS
mSK1a	1	MEPECPRGLLPRPCRVL W LNPRGGKGKALQLE FRSHVQ Q FL EEAEIT
mSK1b	1	MWWCCVL.FVVECPRGLLPRPCRVL W LNPRGGKGKALQLE FRSHVQ Q FL EEAEIT
Yeast LCB4	206	NISSGTVEEILEKSYENSKRNRSLV W LNPRGGKGKALQLE FRSHVQ Q FL EEAEIT
Yeast LCB5	252	DL...VEEILKRSYKNTRENKSIFW W LNPRGGKGKALQLE FRSHVQ Q FL EEAEIT
S. pombe	90	FCEYLLDVAYKG W IKFRRFIVFINPHGGKGKAKKLEM W LNPRGGKGKALQLE FRSHVQ Q FL EEAEIT
C. elegans	63	ENEQLTSVILSRKPPQEQCRGNLL W FINPNNSGTG W KSLET W ANTVGP W EDKSLIR
Consensus		*...R...*V.*NP.GGKGKA.*F.*...P.L...*
hSK	112	SGNA LAASLNHYAGYEQVTNEDLLTNC T LLCRR...LLSP W NLLSLHTASGLRL
mSK1a	111	SGNA LAASVNHYAGYEQVTNEDLLINCT T LLCRR...RLSP W NLLSLHTASGLRL
mSK1b	118	SGNA LAASVNHYAGYEQVTNEDLLINCT T LLCRR...RLSP W NLLSLHTASGLRL
Yeast LCB4	325	SGNA M S IS...CHWTNNPSYAA W CLVKSIETRIDL W CCSQQPSYMNEWPR
Yeast LCB5	368	SGNA MSVS...CHWTNNPSYST W CLIKSIETRIDL W CCSQQPSYAREHPK
S. pombe	203	SGNA F S YN...ATGQLKPALTAD W ELKGRPTSFDL W TFEQ...KQKKA
C. elegans	184	SGNG LLCSV...LSKYGTKMNEKSVMERAEIATSPAKAESVALYSVKTDNQ.SY
Consensus		SGNA*...S...*L.*...*M...

Figure 9A

Figure 9B.

FTLMLTERRN~~H~~ARE. LVRSEELGRW~~D~~ALVVM~~SG~~D~~G~~CLMHEV~~NG~~LMER~~PD~~WETAI. QKPLCSL~~P~~~~A~~
 FKLILTERK~~N~~ARE. LVCAEELGHW~~D~~ALA~~V~~MSGD~~G~~CLMHEV~~NG~~LMER~~PD~~WETAI. QKPLCSL~~P~~~~G~~
 FKLILTERK~~N~~ARE. LVCAEELGHW~~D~~ALA~~V~~MSGD~~G~~CLMHEV~~NG~~LMER~~PD~~WETAI. QKPLCSL~~P~~~~G~~
 IEIAY~~T~~KYARHAID. IAKDLDISKYDTIACAS~~G~~D~~G~~IPYEV~~NG~~LYRR~~PD~~RVDA~~F~~NKLA~~V~~TQL~~P~~~~G~~
 IEVY~~T~~KYPGHAIE. IAREMDIDK~~Y~~DTIACAS~~G~~D~~G~~IPHEV~~NG~~LYQR~~PD~~HVK~~A~~FNNIAITE~~I~~~~P~~~~G~~
 CEVVL~~T~~RRKDHAKS. IAKNLDVG~~S~~Y~~D~~G~~I~~LSV~~G~~D~~G~~CLFHEV~~NG~~LY~~D~~Y~~L~~EA~~F~~. RLPVCMI~~P~~~~G~~
 YEVVV~~T~~TGPN~~H~~ARNVLM~~T~~KADLGK~~F~~NGV~~L~~LS~~G~~D~~G~~GLV~~F~~EAL~~N~~GLC~~R~~EDAFRIFTPLPIGIVE~~S~~~~G~~
 * * . T . . . HA . * . . . * D . * . . . SGD~~G~~ * . . . EV * NGL . . R . D . . . A . . . * . . . * P . G

F~~S~~V~~L~~SLAW~~G~~FIADVD~~L~~ES~~E~~KYRR . L~~G~~EMRF~~T~~L~~G~~TFRLA~~A~~LR~~T~~~~Y~~RGRL
 Y~~S~~V~~L~~LSW~~G~~FVADVD~~L~~ES~~E~~KYRR . L~~G~~EIRF~~T~~VG~~T~~FFRLA~~S~~LR~~I~~~~Y~~QGQL
 Y~~S~~V~~L~~LSW~~G~~FVADVD~~L~~ES~~E~~KYRR . L~~G~~EIRF~~T~~VG~~T~~FFRLA~~S~~LR~~I~~~~Y~~QGQL
 LSFLSQT~~Y~~CVIAESDINTE . FIRWMGPVR~~N~~LG~~V~~A~~F~~N~~I~~IQ~~G~~KK~~Y~~PC~~E~~V~~F~~V~~K~~YAA~~K~~SK~~K~~KELKVHFL
 LSFLSQT~~Y~~GLIAETDINTE . FIRWMGPAREEL~~G~~V~~A~~~~F~~N~~I~~IQ~~G~~KK~~Y~~PC~~E~~IY~~V~~K~~Y~~AA~~K~~SK~~N~~ELKNH~~Y~~L
 Y~~S~~FET~~Y~~T~~AN~~Y~~G~~IIADC~~D~~DIGTE . NWRFMGENRAYL~~G~~FF~~L~~RL~~F~~Q~~K~~PDW~~K~~CS~~I~~EMDV~~V~~SSDR~~T~~EIKH~~M~~
 ASFESIGW~~G~~LMAD~~D~~DID~~S~~EKWRK~~S~~LG~~H~~H~~R~~FTVMGF~~I~~RS~~C~~NL~~R~~~~S~~~~Y~~KGRL
 S . L.S . * * G . * A * . D * * E . * R . * G . . RF . * G . . . * . Y . . . *

Figure 9C

Figure 9D

Figure 10A

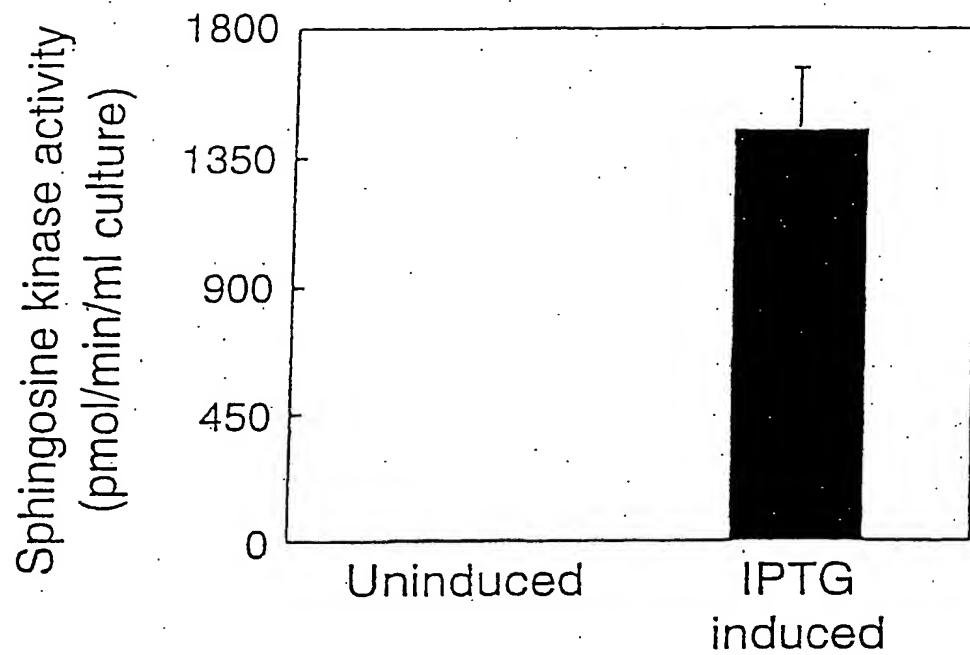


Figure 10B

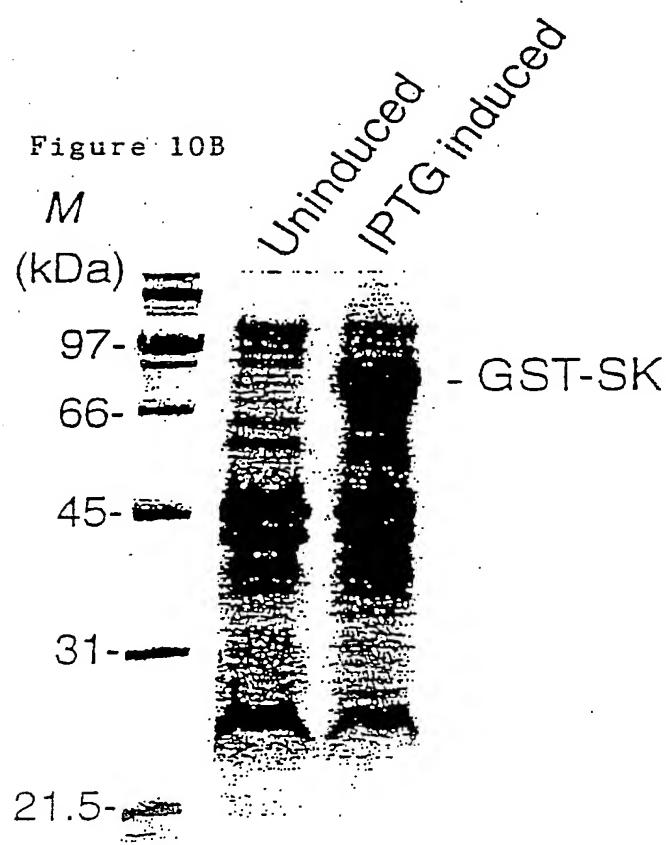
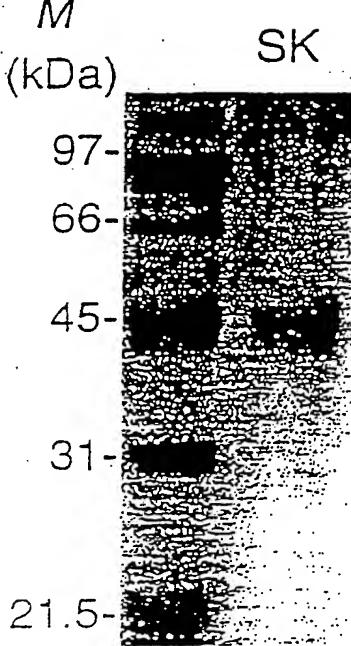


Figure 10C



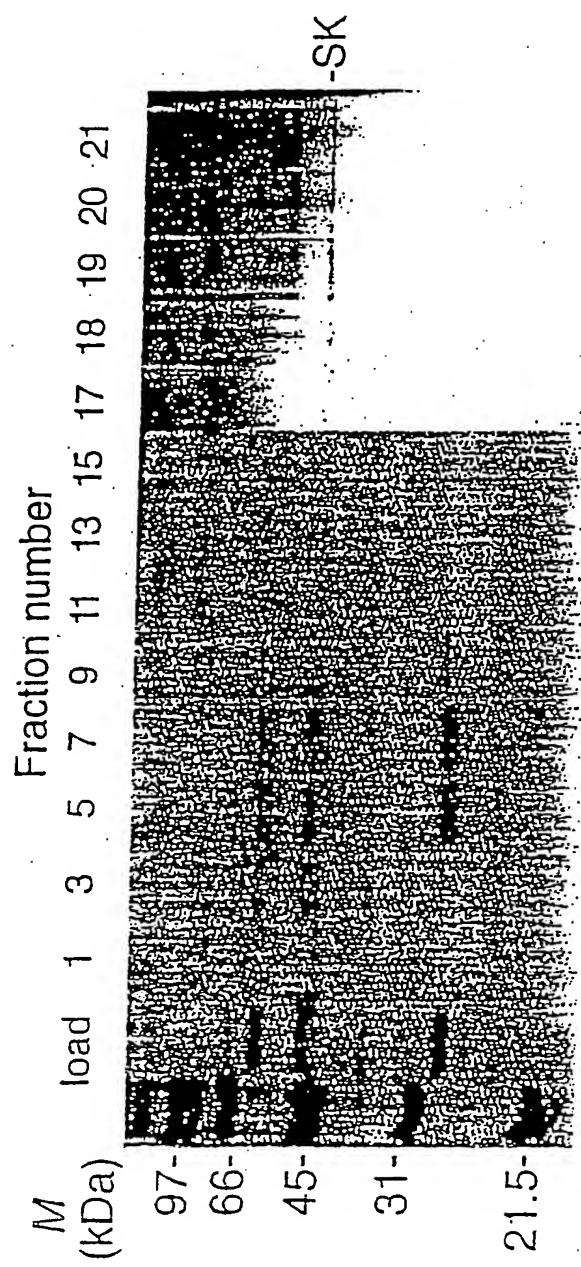


Figure 11A

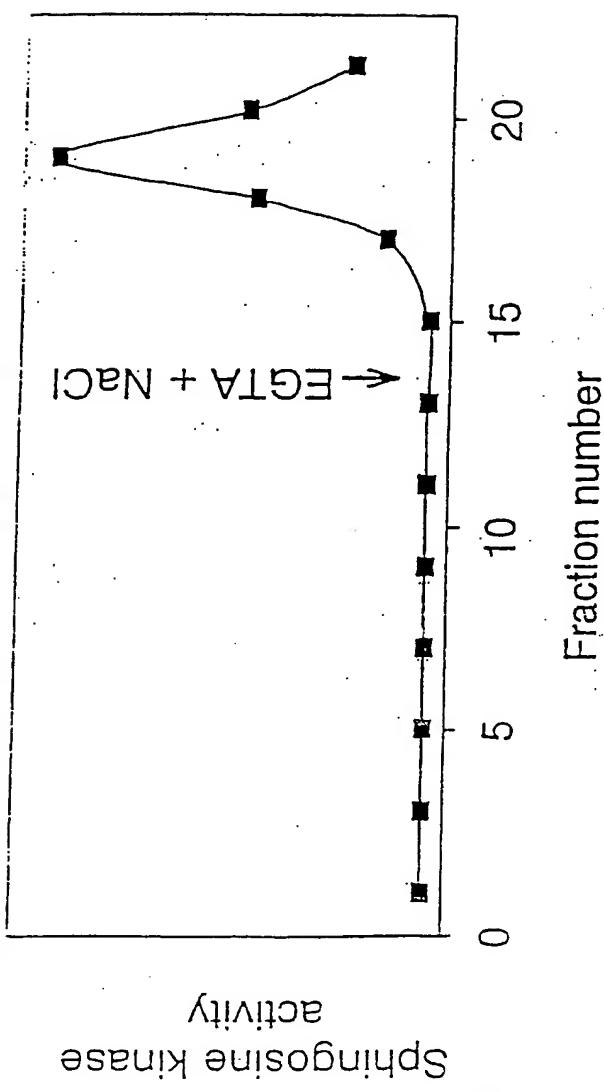


Figure 11B

Figure 12A

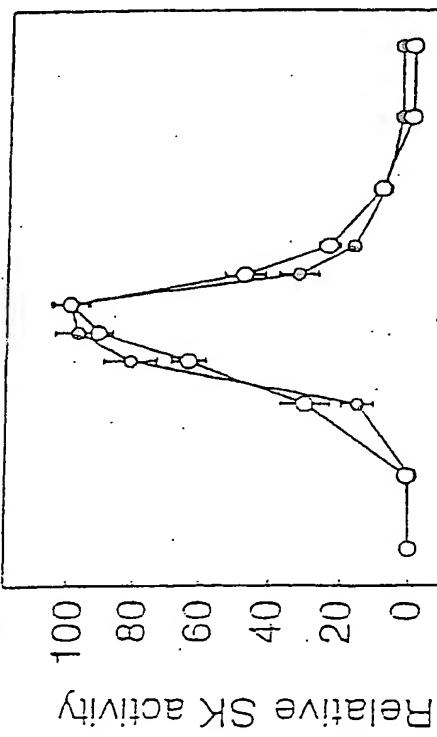
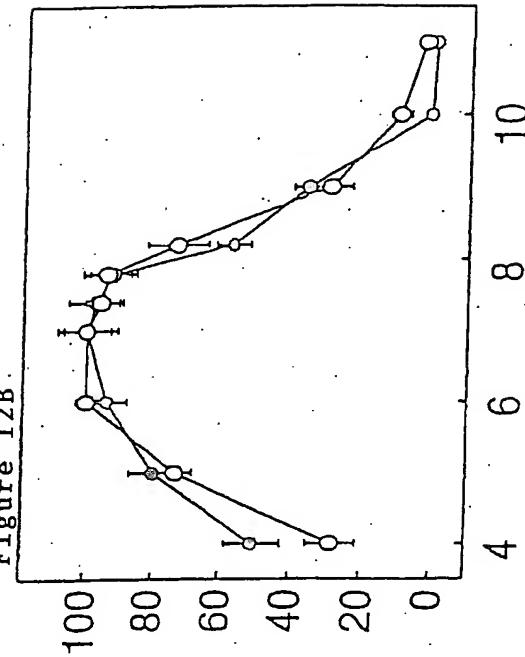


Figure 12B



Relative SK activity

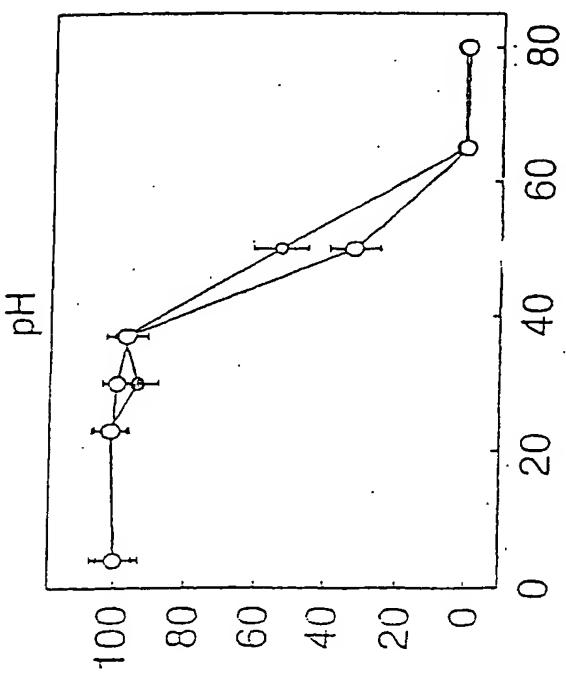


Figure 12C

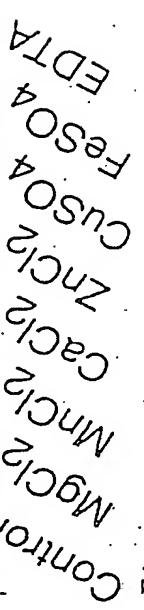
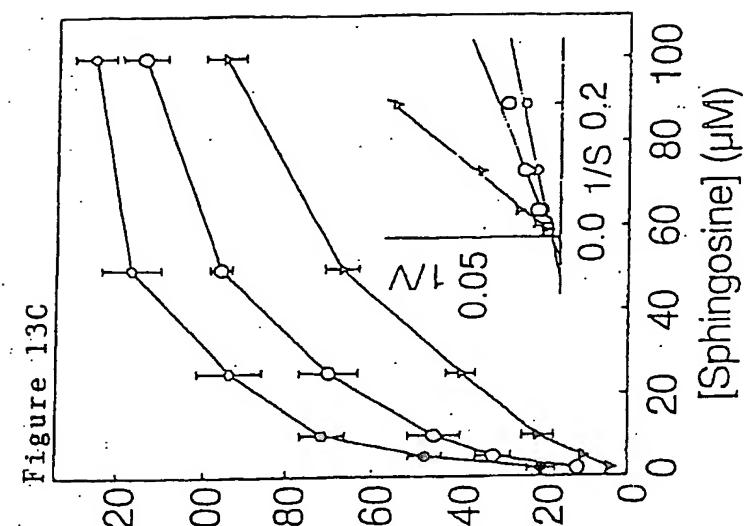
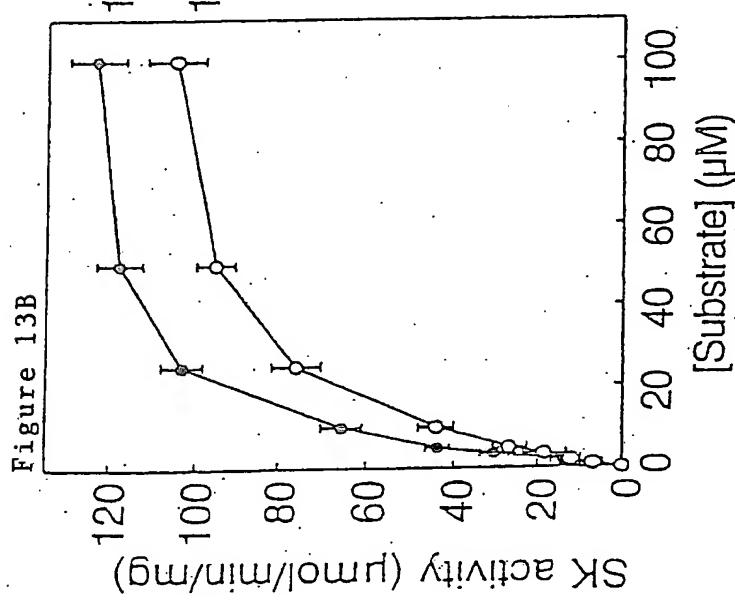
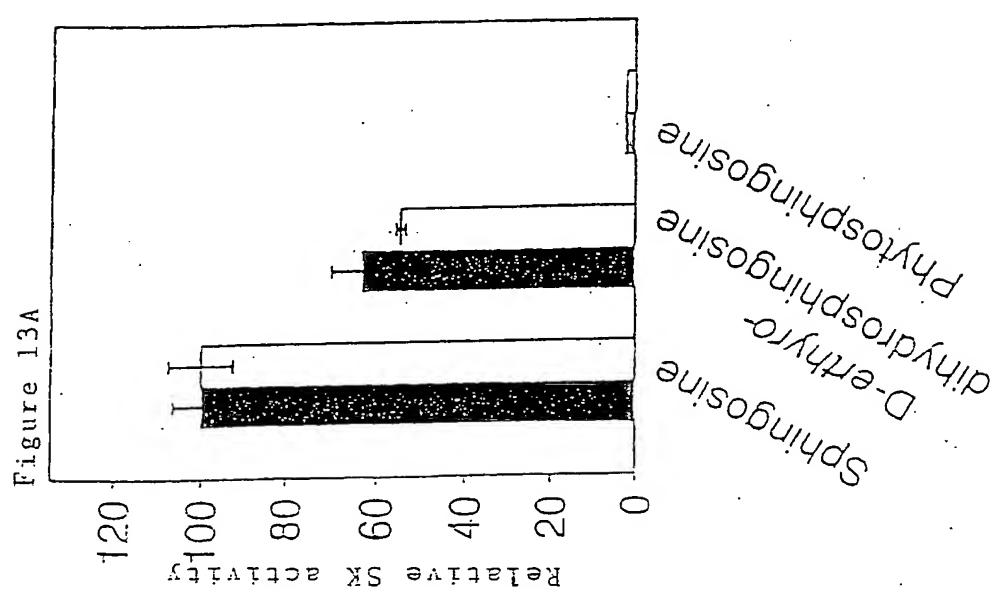


Figure 12D



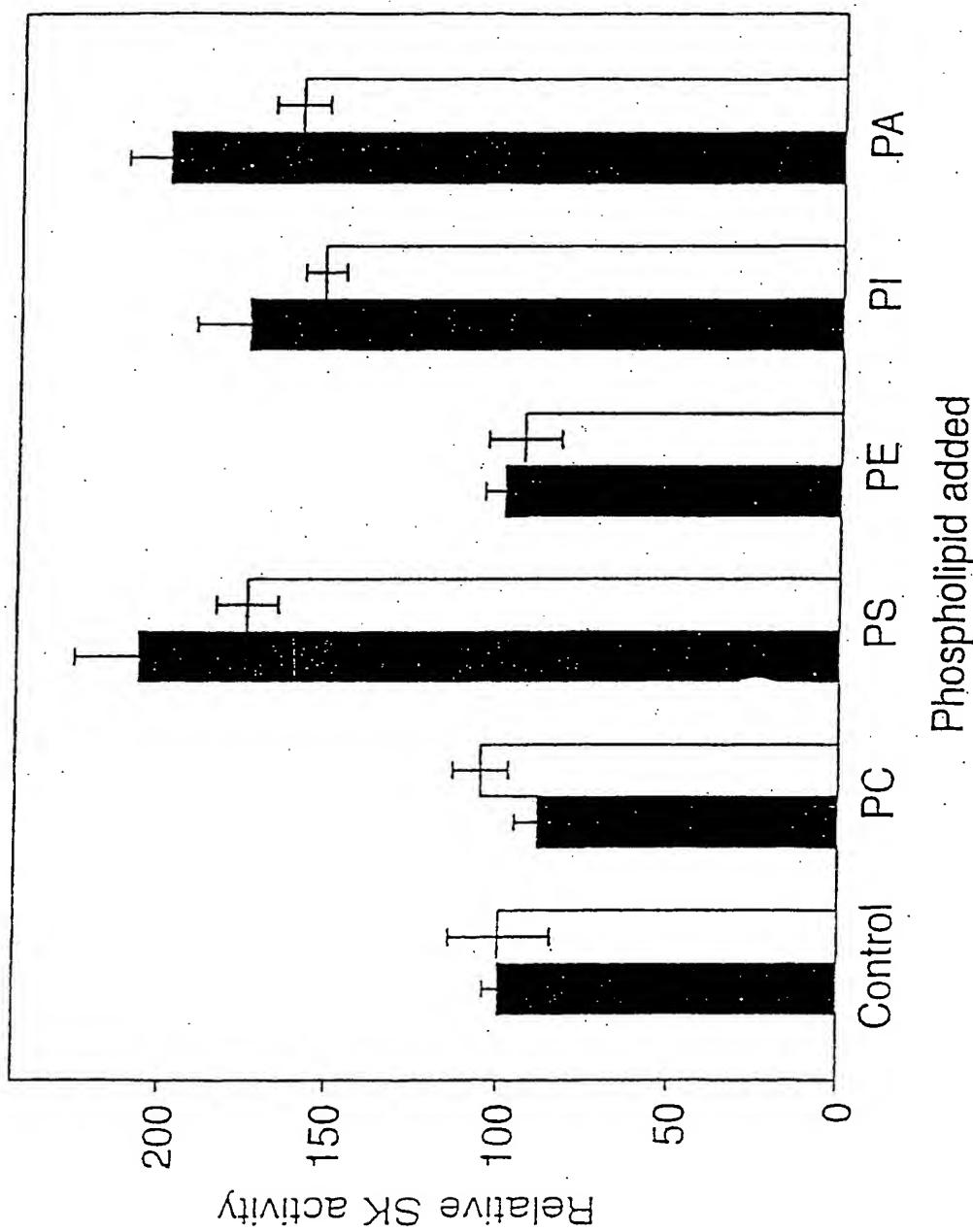


Figure 14